

PCT

RAW SEQUENCE LISTING

DATE: 12/17/2004

PATENT APPLICATION: US/10/516,587

TIME: 14:58:46

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\12172004\J516587.raw

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4 <110> APPLICANT: KANEKA CORPORATION
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- 6 <120> TITLE OF INVENTION: NOVEL ACYLASE GENE
- 8 <130> FILE REFERENCE: T719./ACYL-1
- C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/516,587
- C--> 10 <141> CURRENT FILING DATE: 2004-12-03
 - 10 <150> PRIOR APPLICATION NUMBER: JP P2002-165722
 - 11 <151> PRIOR FILING DATE: 2002-06-06
 - 13 <160> NUMBER OF SEQ ID NOS: 15

ERRORED SEQUENCES

470 <210> SEQ ID NO: 15

471 <211> LENGTH: 26

472 <212> TYPE: DNA

473 <213> ORGANISM: Artificial Sequence

475 <220> FEATURE:

476 <223> OTHER INFORMATION: Description of Artificial Sequence: MT-218 primer

478 <400> SEQUENCE: 15

479 ggaagaagct tgtgcagcac ccggcc

26

Corrected Diskette Needed

Dees Not Comply

E--> 480/1 E--> 482 1/13

E-->484/2/13E--> 486 3/13

E--> 488 4/13

E--> 490 5/13

E--> 492 6/13

E--> 494 7/13

E--> 496 8/13

 $E--> 498 \sqrt{9/13}$

E--> 500 10/13

E--> 502 /11/13

E--> 504/12/13

E--> 506 13/13

VERIFICATION SUMMARY

DATE: 12/17/2004

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:480 M:254 E: No. of Bases conflict, this line has no nucleotides; M:254 Repeated in SeqNo=15 L:482 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:484 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:486 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:488 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:490 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:492 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:494 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:496 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:498 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:500 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3 L:502 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3 L:504 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3 L:506 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3 L:506 M:252 E: No. of Seq. differs, <211> LENGTH: Input:26 Found:39 SEQ:15